

PCT

SAK

## RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/647,019

TIME: 16:23:00

Input Set : A:\12525-407001.TXT

Output Set: N:\CRF3\07032001\I647019.raw

4 <110> APPLICANT: Harvey, Richard P.  
5 Palmer, Stephen J.  
6 Rosenthal, Nadia A.  
7 Musaro, Antonio  
9 <120> TITLE OF INVENTION: NOVEL MOLECULES EXPRESSED DURING MUSCLE  
10 DEVELOPMENT AND GENETIC SEQUENCES ENCODING THE SAME  
13 <130> FILE REFERENCE: 12525-407001  
15 <140> CURRENT APPLICATION NUMBER: 09/647,019  
C--> 16 <141> CURRENT FILING DATE: 2001-06-13  
18 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00220  
19 <151> PRIOR FILING DATE: 1999-03-26  
21 <150> PRIOR APPLICATION NUMBER: AU PP2634/98  
22 <151> PRIOR FILING DATE: 1998-03-27  
24 <160> NUMBER OF SEQ ID NOS: 20  
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
28 <210> SEQ ID NO: 1  
29 <211> LENGTH: 778  
30 <212> TYPE: DNA  
31 <213> ORGANISM: Mus musculus  
33 <220> FEATURE:  
34 <221> NAME/KEY: CDS  
35 <222> LOCATION: (199)...(453)  
37 <400> SEQUENCE: 1

38 gctctcagga ctggagagag acagagcact ccagctatctt cagccacatg aaaagcactg	60
39 gaattgagat ccccgctcag aggacaccgg gagttccttc tctcctgtaa agcgcttttt	120
40 gtgtttttgc acctggccgc ctgggactgt cctcaggcag taaaccaatc cagagagcag	180
41 ggctaagacc ttgtgaat atg tcg aag cag cca att tcc aac gtc aga gcc	231
42 Met Ser Lys Gln Pro Ile Ser Asn Val Arg Ala	
43 1 5 10	
45 atc cag gcg aat atc aat att cca atg gga gcc ttt cgt ccg gga gct	279
46 Ile Gln Ala Asn Ile Asn Ile Pro Met Gly Ala Phe Arg Pro Gly Ala	
47 15 20 25	
49 ggg cag cct ccc aga agg aaa gag agt act cct gaa act gag gag gga	327
50 Gly Gln Pro Pro Arg Arg Lys Glu Ser Thr Pro Glu Thr Glu Glu Gly	
51 30 35 40	
53 gct cct acc acc tca gag gaa aag aag cca att cct gga atg aag aaa	375
54 Ala Pro Thr Thr Ser Glu Glu Lys Lys Pro Ile Pro Gly Met Lys Lys	
55 45 50 55	
57 ttt cca gga cct gtt gtc aac ttg tct gag atc caa aat gtt aaa agt	423
58 Phe Pro Gly Pro Val Val Asn Leu Ser Glu Ile Gln Asn Val Lys Ser	
59 60 65 70 75	
61 gaa ctg aaa ttt gtc ccc aaa ggt gaa cag tagtcgaaag gacacaaaag	473
62 Glu Leu Lys Phe Val Pro Lys Gly Glu Gln	
63 80 85	
65 ttacacattgg atgcttagaa tcaggagatg catttcgctt acgtgttttt ccaagggaga	533
66 aaaaacaatg ggttgaaata aacaacttcc tgaacatttt atacatttgt atgatgatca	593
67 caaacctcct gaatgcccaa gactctagca aaaatatcct gttgtacat ttatatttct	653

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68 tccttttact tggttgcatt tctcacttta gctacatttt tggcaccttg tagagcaaat 713
69 cagcacacga atttacaacc tgggaagtgt ggttttgagg agagatgtga tttttatgaa 773
70 gggggg 778
72 <210> SEQ ID NO: 2
73 <211> LENGTH: 85
74 <212> TYPE: PRT
75 <213> ORGANISM: Mus musculus
77 <400> SEQUENCE: 2
78 Met Ser Lys Gln Pro Ile Ser Asn Val Arg Ala Ile Gln Ala Asn Ile
79 1 5 10 15
80 Asn Ile Pro Met Gly Ala Phe Arg Pro Gly Ala Gly Gln Pro Pro Arg
81 20 25 30
82 Arg Lys Glu Ser Thr Pro Glu Thr Glu Glu Gly Ala Pro Thr Thr Ser
83 35 40 45
84 Glu Glu Lys Lys Pro Ile Pro Gly Met Lys Lys Phe Pro Gly Pro Val
85 50 55 60
86 Val Asn Leu Ser Glu Ile Gln Asn Val Lys Ser Glu Leu Lys Phe Val
87 65 70 75 80
88 Pro Lys Gly Glu Gln
89 85
91 <210> SEQ ID NO: 3
92 <211> LENGTH: 887
93 <212> TYPE: DNA
94 <213> ORGANISM: Homo sapiens
96 <220> FEATURE:
97 <221> NAME/KEY: CDS
98 <222> LOCATION: (185)...(448)
100 <400> SEQUENCE: 3
101 ggttctcaat accgggagag gcacagagct atttcagcca catgaaaagc atcggaattg 60
102 agatcgagcgc tcagaggaca ccgggcgcct cttccacctt ccaaggagct ttgtattctt 120
103 gcactctggct gcctgggact tcccttaggc agtaaacaaa tacataaagc agggataaga 180
104 ctgc atg aat atg tcg aaa cag cca gtt tcc aat gtt aga gcc atc cag 229
105 Met Asn Met Ser Lys Gln Pro Val Ser Asn Val Arg Ala Ile Gln
106 1 5 10 15
108 gca aat atc aat att cca atg gga gcc ttt cgg cca gga gca ggt caa 277
109 Ala Asn Ile Asn Ile Pro Met Gly Ala Phe Arg Pro Gly Ala Gly Gln
110 20 25 30
112 ccc ccc aga aga aaa gaa tgt act cct gaa gtg gag gag ggt gtt cct 325
113 Pro Pro Arg Arg Lys Glu Cys Thr Pro Glu Val Glu Glu Gly Val Pro
114 35 40 45
116 ccc acc tcg gat gag gag aag aag cca att cca gga gcg aag aaa ctt 373
117 Pro Thr Ser Asp Glu Glu Lys Lys Pro Ile Pro Gly Ala Lys Lys Leu
118 50 55 60
120 cca gga cct gca gtc aat cta tcg gaa atc cag aat att aaa agt gaa 421
121 Pro Gly Pro Ala Val Asn Leu Ser Glu Ile Gln Asn Ile Lys Ser Glu
122 65 70 75
124 cta aaa tat gtc ccc aaa gct gaa cag tagtaggaag aaaaaaggat 468
125 Leu Lys Tyr Val Pro Lys Ala Glu Gln
126 80 85

```

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128 tgatgtgaag aaataaagag gcagaagatg gattcaatag ctcaactaaa ttttatatat 528
129 ttgtatgatg attgtgaacc tcctgaatgc ctgagactct agcagaaatg gcctgtttgt 588
130 acattttatat ctcttccttc tagttggctg tatttcttac tttatcttca tttttggcac 648
131 ctcacagaac aaattagccc ataaattcaa cacctggagg gtgtgggtttt gaggagggat 708
132 atgattttat ggagaatgat atggcaatgt gcctaacgat tttgatgaaa agtttcccaa 768
133 gctacttcct acagtattttt ggtcaatatt tggaatgcgt tttagttctt caccttttaa 828
134 attatgtcac taaactttgt atgagttcaa ataaatattt gactaaatgt aaaatgtga 887

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136 &lt;210&gt; SEQ ID NO: 4

137 &lt;211&gt; LENGTH: 88

138 &lt;212&gt; TYPE: PRT

139 &lt;213&gt; ORGANISM: Homo sapiens

141 &lt;400&gt; SEQUENCE: 4

142 Met Asn Met Ser Lys Gln Pro Val Ser Asn Val Arg Ala Ile Gln Ala

143 1 5 10 15

144 Asn Ile Asn Ile Pro Met Gly Ala Phe Arg Pro Gly Ala Gly Gln Pro

145 20 25 30

146 Pro Arg Arg Lys Glu Cys Thr Pro Glu Val Glu Glu Gly Val Pro Pro

147 35 40 45

148 Thr Ser Asp Glu Glu Lys Lys Pro Ile Pro Gly Ala Lys Lys Leu Pro

149 50 55 60

150 Gly Pro Ala Val Asn Leu Ser Glu Ile Gln Asn Ile Lys Ser Glu Leu

151 65 70 75 80

152 Lys Tyr Val Pro Lys Ala Glu Gln

153 85

155 &lt;210&gt; SEQ ID NO: 5

156 &lt;211&gt; LENGTH: 75

157 &lt;212&gt; TYPE: PRT

158 &lt;213&gt; ORGANISM: Xenopus laevis

160 &lt;400&gt; SEQUENCE: 5

161 Met Ser Lys Gln Pro Ala Ser Asn Ile Arg Ser Ile Gln Ala Asn Ile

162 1 5 10 15

163 Asn Ile Pro Met Gly Ala Phe Arg Pro Gly Ala Gly Gln Pro Pro Lys

164 20 25 30

165 Arg Lys Glu Phe Ser Thr Glu Glu Glu Gln His Val Pro Thr Pro Glu

166 35 40 45

167 Ser Glu Glu Lys Ser Glu Glu Lys Lys Pro Ile Pro Gly Ala Val Lys

168 50 55 60

169 Leu Pro Gly Pro Ala Phe Asn Leu Ser Glu Thr

170 65 70 75

172 &lt;210&gt; SEQ ID NO: 6

173 &lt;211&gt; LENGTH: 172

174 &lt;212&gt; TYPE: DNA

175 &lt;213&gt; ORGANISM: Homo sapiens

177 &lt;400&gt; SEQUENCE: 6

178 ggttctcaat accgggagag gcacagagct atttcagcca catgaaaagc atcggaattg 60

179 agatcgcagc tcagaggaca ccgggcgccc ctccacatt ccaaggagct ttgtattctt 120

180 gcatctggct gcctgggact tcccttaggc agtaaacaaa tacataaagc ag 172

182 &lt;210&gt; SEQ ID NO: 7

183 &lt;211&gt; LENGTH: 57

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184 <212> TYPE: DNA
185 <213> ORGANISM: Homo sapiens
187 <400> SEQUENCE: 7
188 ggataagact gcatgaatat gtcgaaacag ccagtttcca atgtagagc catccag      57
190 <210> SEQ ID NO: 8
191 <211> LENGTH: 87
192 <212> TYPE: DNA
193 <213> ORGANISM: Homo sapiens
195 <400> SEQUENCE: 8
196 gcaaatatca atattccaat gggagccttt cggccaggag caggtcaacc cccagaaga      60
197 aaagaatgta ctccctgaagt ggaggag      87
199 <210> SEQ ID NO: 9
200 <211> LENGTH: 149
201 <212> TYPE: DNA
202 <213> ORGANISM: Homo sapiens
204 <400> SEQUENCE: 9
205 ggtgttcctc ccacctcgga tgaggagaag aagccaattc caggagcgaa gaaacttcca      60
206 ggacctgcag tcaatctatc ggaaatccag aatattaaaa gtgaactaaa atatgtcccc      120
207 aaagctgaac agtagtagga agaaaaaag      149
209 <210> SEQ ID NO: 10
210 <211> LENGTH: 422
211 <212> TYPE: DNA
212 <213> ORGANISM: Homo sapiens
214 <400> SEQUENCE: 10
215 gattgatgtg aagaaataaa gaggcagaag atggattcaa tagctcacta aaattttata      60
216 tatttgtatg atgattgtga acctcctgaa tgcctgagac tctagcagaa atggcctggt      120
217 tgtacattta tatctcttcc ttctagttgg ctgtatttct tactttatct tcatttttgg      180
218 cacctcacag aacaaattag cccataaatt caacacctgg aggggtgtgg tttgaggagg      240
219 gatatgattt tatggagaat gatatggcaa tgtgcctaac gattttgatg aaaagtttcc      300
220 caagctactt cctacagtat tttggtcaat atttggaatg cgttttagtt cttcaccttt      360
221 taaattatgt cactaaactt tgtatgagtt caaataaata tttgactaaa tgtaaaatgt      420
222 ga      422
224 <210> SEQ ID NO: 11
225 <211> LENGTH: 40
226 <212> TYPE: PRT
227 <213> ORGANISM: Patinopecten sp.
229 <400> SEQUENCE: 11
230 Ser Val Ile Gln Arg Asn Ile Arg Lys Trp Val Leu Arg Leu Asn Trp
231 1 5 10 15
232 Gln Trp Trp Lys Leu Tyr Ser Lys Val Lys Pro Leu Leu Ser Ile Ala
233 20 25 30
234 Arg Gln Glu Glu Glu Met Lys Glu
235 35 40
237 <210> SEQ ID NO: 12
238 <211> LENGTH: 40
239 <212> TYPE: PRT
240 <213> ORGANISM: Rattus norvegicus
242 <400> SEQUENCE: 12
243 Leu Val Ile Gln Trp Asn Ile Arg Ala Phe Met Gly Val Lys Asn Trp

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244 1          5          10          15
245 Pro Trp Met Lys Leu Tyr Phe Lys Ile Lys Pro Leu Leu Lys Ser Ala
246          20          25          30
247 Glu Thr Glu Lys Glu Met Ala Asn
248          35          40
250 <210> SEQ ID NO: 13
251 <211> LENGTH: 40
252 <212> TYPE: PRT
253 <213> ORGANISM: Rattus norvegicus
255 <400> SEQUENCE: 13
256 Phe Cys Ile Gln Tyr Asn Ile Arg Ala Phe Met Asn Val Lys His Trp
257 1          5          10          15
258 Pro Trp Met Lys Leu Phe Phe Lys Ile Lys Pro Leu Leu Lys Ser Ala
259          20          25          30
260 Glu Thr Glu Lys Glu Met Ala Thr
261          35          40
263 <210> SEQ ID NO: 14
264 <211> LENGTH: 40
265 <212> TYPE: PRT
266 <213> ORGANISM: Homo sapiens
268 <400> SEQUENCE: 14
269 Glu Asn Asn Val Met Asn Ile Arg Gln Phe Asn Cys Ser Pro His Pro
270 1          5          10          15
271 Tyr Trp Leu Pro Asn Phe Met Asp Val Phe Thr Trp Ser Leu Pro Phe
272          20          25          30
273 Val Gly Glu Lys Arg Val Thr Glu
274          35          40
276 <210> SEQ ID NO: 15
277 <211> LENGTH: 6
278 <212> TYPE: PRT
279 <213> ORGANISM: Artificial Sequence
281 <220> FEATURE:
282 <223> OTHER INFORMATION: consensus sequence
284 <221> NAME/KEY: VARIANT
285 <222> LOCATION: (1)...(6)
286 <223> OTHER INFORMATION: Xaa = Any Amino Acid
288 <400> SEQUENCE: 15
W--> 289 Ile Gln Xaa Xaa Ile Arg
290 1          5
292 <210> SEQ ID NO: 16
293 <211> LENGTH: 42
294 <212> TYPE: PRT
295 <213> ORGANISM: Homo sapiens
297 <400> SEQUENCE: 16
298 Ser Glu Glu Asp Gly Phe Asp Gly Ala Thr Ala Ala Ala Arg Lys Glu
299 1          5          10          15
300 Val Ile Arg Trp Lys Ile Arg Ala Ile Gly Lys Met Ala Arg Val Phe
301          20          25          30
302 Ser Val Leu Arg Glu Glu Ser Glu Ser Val

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**VERIFICATION SUMMARY**

**PATENT APPLICATION:** US/09/647,019

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L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15